/FW/6

CRF Errors Edited by the STIC Systems Branch

		,
Number:	08/9/6,/40	CRF Edit Date: 10/8
	nucleic acid/amino acid numb	pers/text in cases where the sequen
Corrected	the SEQ ID NO. Sequence no	umbers edited were:
Inserted or NO's edit		at the end of a nucleic line. SEQ II
Deleted: _	invalid beginning/end-of-f	ile text ; page numbers
Inserted m	nandatory headings/numeric i	dentifiers, specifically:
Moved res	sponses to same line as headin	g/numeric identifier, specifically:
	1	g/numeric identifier, specifically: 1 spelling of Beetle



IFW16

RAW SEQUENCE LISTING DATE: 10/08/2004
PATENT APPLICATION: US/08/916,140 TIME: 18:19:29

Input Set : A:\PTO.AMC.TXT

```
4 <110 > APPLICANT: Scott Mathew P.
           Goodrich, Lisa V.
   5
           Johnson, Ronald L.
           Epstein, Ervin Jr.
   9 <120> TITLE OF INVENTION: PATCHED GENES AND USES RELATED THERETO
   12 <130> FILE REFERENCE: CIBT-P04-203
   14 <140> CURRENT APPLICATION NUMBER: US 08/916,140
   15 <141> CURRENT FILING DATE: 1997-08-21
  17 <150> PRIOR APPLICATION NUMBER: US 08/656,055
   18 <151> PRIOR FILING DATE: 1996-05-31
   20 <150> PRIOR APPLICATION NUMBER: US 08/540,406
   21 <151> PRIOR FILING DATE: 1995-10-06
   23 <150> PRIOR APPLICATION NUMBER: US 08/319,745
   24 <151> PRIOR FILING DATE: 1994-10-07
   26 <160> NUMBER OF SEQ ID NOS: 64
   28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
   30 <210> SEQ ID NO: 1
   31 <211> LENGTH: 736
   32 <212> TYPE: DNA
   33 <213> ORGANISM: Beetle
   35 <220> FEATURE:
   37 <221> NAME/KEY: misc feature
   38 <222> LOCATION: 4, 5, 7, 8, 10, 11, 23, 34, 35, 36, 39, 40, 41, 45, 51, 52,
            57, 61, 71, 75, 77, 87, 88, 89, 91, 92, 96, 97, 100, 104,
   39
   40
            106, 109, 111, 113, 117, 120, 126, 149, 151, 153, 154, 157,
           178, 187, 189, 191, 211, 214, 310, 608, 704, 708, 712
   41
   42 <223> OTHER INFORMATION: n = A, T, C or G
   44 <221> NAME/KEY: misc_feature
   45 <222> LOCATION: 714, 729, 732
   46 <223 > OTHER INFORMATION: n = A, T, C \text{ or } G
-> 48 <400> 1
-> 49 aacnncnntn natggcaccc ccncccaacc tttnnnccnn ntaancaaaa nnccccnttt 60
   50 natacccct ntaananttt tccaccnnnc nnaaannccn ctgnanacna ngnaaanccn 120
   51 tttttnaacc cccccaccc ggaattccna ntnnccnccc ccaaattaca actccagncc 180
   52 aaaattnana naattggtcc taacctaacc natngttgtt acggtttccc cccccaaata 240
   53 catgcactgg cccgaacact tgatcgttgc cgttccaata agaataaatc tggtcatatt 300
   54 aaacaaqccn aaaqctttac aaactqttgt acaattaatg ggcgaacacg aactgttcga 360
   55 attotggtot ggacattaca aagtgcacca categgatgg aaccaggaga aggccacaac 420
   56 cgtactgaac gcctggcaga agaagttcgc acaggttggt ggttggcgca aggagtagag 480
   57 tgaatggtgg taatttttgg ttgttccagg aggtggatcg tctgacgaag agcaagaagt 540
   58 cgtcgaatta catcttcgtg acgttctcca ccgccaattt gaacaagatg ttgaaggagg 600
   59 cgtcgaanac ggacgtggtg aagctggggg tggtgctggg ggtggcggcg gtgtacgggt 660
   60 gggtggccca gtcggggctg gctgccttgg gagtgctggt cttngcgngc tncnattcgc 720
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RAW SEQUENCE LISTINGPATENT APPLICATION: **US/08/916,140**DATE: 10/08/2004

TIME: 18:19:29

Input Set : A:\PTO.AMC.TXT

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736
    61 cctatagtna gncgta
    63 <210> SEQ ID NO: 2
    64 <211> LENGTH: 107
    65 <212> TYPE: PRT
    66 <213> ORGANISM: Beetle
    68 <220> FEATURE:
    70 <221> NAME/KEY: VARIANT
    71 <222> LOCATION: 1, 9, 12, 13, 14, 20
    72 <223> OTHER INFORMATION: Xaa = Any Amino Acid
    74 <400> SEQUENCE: 2
W--> 75 Xaa Pro Pro Pro Asn Tyr Asn Ser Xaa Pro Lys Xaa Xaa Xaa Leu Val
    77 Leu Thr Pro Xaa Val Val Thr Val Ser Pro Pro Lys Tyr Met His Trp
    79 Pro Glu His Leu Ile Val Ala Val Pro Ile Arg Ile Asn Leu Val Ile
    80
    81 Leu Asn Lys Pro Lys Ala Leu Gln Thr Val Val Gln Leu Met Gly Glu
                                55
    83 His Glu Leu Phe Glu Phe Trp Ser Gly His Tyr Lys Val His His Ile
    85 Gly Trp Asn Gln Glu Lys Ala Thr Thr Val Leu Asn Ala Trp Gln Lys
                        85
    87 Lys Phe Ala Gln Val Gly Gly Trp Arg Lys Glu
    88
                    100
    91 <210> SEQ ID NO: 3
    92 <211> LENGTH: 5187
    93 <212> TYPE: DNA
    94 <213> ORGANISM: Butterfly
    96 <400> SEQUENCE: 3
    97 ggqtctqtca cccqqaqccq gagtccccgg cggccagcag cgtcctcgcg agccgagcgc 60
    98 ccaggegege eeggageeg eggeggegge ggcaacatgg ceteggetgg taaegeegee 120
     99 ggggccctgg gcaggcaggc cggcggggg aggcgcagac ggaccggggg accgcaccgc 180
    100 gccgcgcgg accgggacta tctgcaccgg cccagctact gcgacgccgc cttcgctctg 240
    101 gaqcaqattt ccaaggggaa ggctactggc cggaaagcgc cgctgtggct gagagcgaag 300
    102 tttcaqaqac tcttatttaa actgggttgt tacattcaaa agaactgcgg caagtttttg 360
    103 qttqtqqqtc tcctcatatt tqqqqccttc qctqtqgqat taaaggcagc taatctcgag 420
    104 accaacgtgg aggagetgtg ggtggaagtt ggtggacgag tgagtcgaga attaaattat 480
    105 accepteaga agataggaga agaggetatg tttaateete aaeteatgat acagaeteea 540
    106 aaagaagaag gcgctaatgt tctgaccaca gaggctctcc tgcaacacct ggactcagca 600
    107 ctccaqqcca qtcqtqtqca cqtctacatq tataacaggc aatggaagtt ggaacatttg 660
    108 tgctacaaat caggggaact tatcacggag acaggttaca tggatcagat aatagaatac 720
    109 ctttaccctt gcttaatcat tacacctttg gactgcttct gggaaggggc aaagctacag 780
    110 tccgggacag catacctcct aggtaagcct cctttacggt ggacaaactt tgaccccttg 840
    111 gaatteetag aagagttaaa gaaaataaac taccaagtgg acagetggga ggaaatgetg 900
    112 aataaagccg aagttggcca tgggtacatg gaccggcctt gcctcaaccc agccgaccca 960
    113 gattgccctg ccacagcccc taacaaaaat tcaaccaaac ctcttgatgt ggcccttgtt 1020
    114 ttgaatggtg gatgtcaagg tttatccagg aagtatatgc attggcagga ggagttgatt 1080
    115 gtgggtggta ccgtcaagaa tgccactgga aaacttgtca gcgctcacgc cctgcaaacc 1140
    116 atgttccagt taatgactcc caagcaaatg tatgaacact tcaggggcta cgactatgtc 1200
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/916,140

DATE: 10/08/2004 TIME: 18:19:29

Input Set : A:\PTO.AMC.TXT

						_	
			agacagggca				
			aagtgtcgcc				
			catcctaaaa				
			gcttgcctat				
			ggggctggct				
			cttgattggc				
			tgttggtgtg				
			gaggattcca				
			cctcacctcc				
			cctgcgagcg				
			gctcattttt				
			tattttctgc				
			ctacacagag				
			cttcgcccac				
			ccctcacacg				
			tgttaccgtc				
133	gagagcacca	gctctaccag	ggacctgctc	tcccagttct	cagactccag	cctccactgc	2220
			gtggacactc				
			caaggttgtg				
			ccgagtgaga				
137	cgggaaacca	gagaatatga	cttcatagct	gcccagttca	agtacttctc	tttctacaac	2460
			agcagactac				
			gaagtatgtc				
			agactggctt				
			gccaaacaat				
			gactggcagc				
143	actaaacagc	gtctggtaga	cgcagatggc	atcattaatc	cgagcgcttt	ctacatctac	2820
			cgaccctgta				
			ccatgacaaa				
			cgagtacgct				
			agccatagaa				
			ccccaatggc				
149	agcctgcgcc	actggctgct	gctatccatc	agcgtggtgc	tggcctgcac	gtttctagtg	3180
			cccctggacg				
			catgatgggc				
			tgttggcatc				
			ggacaagaac				
			tgctgtgtcc				
			cagatacttc				
156	ggggttctca	atggactggt	tctgctgcct	gtcctcttat	ccttctttgg	accgtgtcct	3600
157	gaggtgtctc	cagccaatgg	cctaaaccga	ctgcccactc	cttcgcctga	gccgcctcca	3660
158	agtgtcgtcc	ggtttgccgt	gcctcctggt	cacacgaaca	atgggtctga	ttcctccgac	3720
			cacggtgtct				
			ccctgcccac				
			ggtccatccg				
			ctctggctcc				
			aggcttgcgg				
			gcattctggc				
			teggaaceca				

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/08/916,140**DATE: 10/08/2004 TIME: 18:19:29

Input Set : A:\PTO.AMC.TXT

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166 agetaetgee ageceateae eactgtgaeg gettetgett eggtgaetgt tgetgtgeat 4200
167 cccccgcctg gacctgggcg caacccccga ggggggccct gtccaggcta tgagagctac 4260
168 cctgagactg atcacggggt atttgaggat cctcatgtgc cttttcatgt caggtgtgag 4320
169 aggagggact caaaggtgga ggtcatagag ctacaggacg tggaatgtga ggagaggccg 4380
170 tgggggagca getecaactg agggtaatta aaatetgaag caaagaggee aaagattgga 4440
171 aageeeegee eecacetett teeagaactg ettgaagaga actgettgga attatgggaa 4500
172 ggcagttcat tgttactgta actgattgta ttattkkgtg aaatatttct ataaatattt 4560
173 aaraggtgta cacatgtaat atacatggaa atgctgtaca gtctatttcc tggggcctct 4620
174 ccacteetge eecagagtgg ggagaceaca ggggeeettt eecetgtgta cattggtete 4680
175 tgtgccacaa ccaagcttaa cttagtttta aaaaaaatct cccagcatat gtcgctgctg 4740
176 cttaaatatt gtataattta cttgtataat tctatgcaaa tattgcttat gtaataggat 4800
177 tatttgtaaa ggtttctgtt taaaatattt taaatttgca tatcacaacc ctgtggtagg 4860
178 atgaattgtt actgttaact titgaacacg ctatgcgtgg taattgttta acgagcagac 4920
179 atgaagaaaa caggttaatc ccagtggctt ctctaggggt agttgtatat ggttcgcatg 4980
180 qqtqqatqtq tqtqtqcatq tqactttcca atqtactqta ttqtqqtttq ttqttqttqt 5040
181 tgctgttgtt gttcattttg gtgtttttgg ttgctttgta tgatcttagc tctggcctag 5100
182 gtgggetggg aaggtecagg tetttttetg tegtgatget ggtggaaagg tgaeeceaat 5160
183 catctgtcct attctctggg actattc
                                                                       5187
185 <210> SEQ ID NO: 4
186 <211> LENGTH: 1311
187 <212> TYPE: PRT
188 <213> ORGANISM: Butterfly
190 <220> FEATURE:
192 <221> NAME/KEY: VARIANT
193 <222> LOCATION: 348, 908
194 <223> OTHER INFORMATION: Xaa = Any Amino Acid
196 <400> SEQUENCE: 4
197 Met Val Ala Pro Asp Ser Glu Ala Pro Ser Asn Pro Arg Ile Thr Ala
199 Ala His Glu Ser Pro Cys Ala Thr Glu Ala Arg His Ser Ala Asp Leu
200
201 Tyr Ile Arg Thr Ser Trp Val Asp Ala Ala Leu Ala Leu Ser Glu Leu
203 Glu Lys Gly Asn Ile Glu Gly Gly Arg Thr Ser Leu Trp Ile Arg Ala
                            55
205 Trp Leu Gln Glu Gln Leu Phe Ile Leu Gly Cys Phe Leu Gln Gly Asp
                        70
                                            75
207 Ala Gly Lys Val Leu Phe Val Ala Ile Leu Val Leu Ser Thr Phe Cys
                                        90
209 Val Gly Leu Lys Ser Ala Gln Ile His Thr Arq Val Asp Gln Leu Trp
210
                100
                                    105
211 Val Gln Glu Gly Gly Arg Leu Glu Ala Glu Leu Lys Tyr Thr Ala Gln
212
                                120
                                                     125
213 Ala Leu Gly Glu Ala Asp Ser Ser Thr His Gln Leu Val Ile Gln Thr
                            135
215 Ala Lys Asp Pro Asp Val Ser Leu Leu His Pro Gly Ala Leu Leu Glu
                        150
217 His Leu Lys Val Val His Ala Ala Thr Arg Val Thr Val His Met Tyr
218
                    165
                                        170
```

RAW SEQUENCE LISTING DATE: 10/08/2004
PATENT APPLICATION: US/08/916,140 TIME: 18:19:29

Input Set : A:\PTO.AMC.TXT

	219 220	Asp	Ile	Glu	Trp 180	Arg	Leu	Lys	Asp	Leu 185	Cys	Tyr	Ser	Pro	Ser 190	Ile	Pro
		Asp	Phe	Glu 195		Tyr	His	His	Ile 200	Ģlu	Ser	Ile	Ile	Asp 205	Asn	Val	Ile
		Pro	Cys 210	Ala	Ile	Ile	Thr	Pro 215	Leu	Asp	Cys	Phe	Trp 220	Glu	Gly	Ser	Lys
		Leu 225	Leu	Gly	Pro	Asp	Tyr 230	Pro	Ile	Tyr	Val	Pro 235	His	Leu	Lys	His	Lys 240
	227 228	Leu	Gln	Trp	Thr	His 245	Leu	Asn	Pro	Leu	Glu 250	Val	Val	Glu	Glu	Val 255	Lys
	229 230	Lys	Leu	Lys	Phe 260	Gln	Phe	Pro	Leu	Ser 265	Thr	Ile	Glu	Ala	Tyr 270	Met	Lys
		Arg	Ala	Gly 275	Ile	Thr	Ser	Ala	Tyr 280	Met	Lys	Lys	Pro	Cys 285	Leu	Asp	Pro
		Thr	Asp 290	Pro	His	Cys	Pro	Ala 295	Thr	Ala	Pro	Asn	Lys 300	Lys	Ser	Gly	His
		Ile 305	Pro	Asp	Val	Ala	Ala 310	Glu	Leu	Ser	His	Gly 315	Cys	Tyr	Gly	Phe	Ala 320
		Ala	Ala	Tyr	Met	His	Trp	Pro	Glu	Gln	Leu 330	Ile	Val	Gly	Gly	Ala 335	Thr
>		Arg	Asn	Ser			Ala	Leu	Arg			Arg	Xaa	Leu			Val
	240	_			340					345				_	350	_	
	242	Val		355					360					365			
	244	Tyr	370					375					380				
	246	Leu 385	_		_		390	_				395					400
	248	Thr				405					410					415	
	250	Ser			420	_				425					430		
	252	Asn		435					440					44 5		•	
	254	Leu	450					455					460				
		Ala	Gly	Val	Leu	Leu		Ser	Ile	Thr	Val	Ala 475	Ala	Gly	Leu	Gly	
	256	465 Cys	7 J ¬	I 011	Ton	Clv	470	Dro	Dho	7 cn	Nlα		Sar	Thr	Gln	Tle	480 Val
	258	Cys	на	neu		485		PIO	FILE		490		261	TIII	GIII	495	
		Pro	Phe	Leu	Ala	Leu	Gly	Leu	Gly	Val	Gln	Asp	Met	Phe	Leu	Leu	Thr
	260				500				0	505				_	510		
	262	His		515					520					525			
	264	Gly	530			_		535					540				
		Cys	Asn	Val	Met	Ala		Leu	Ala	Ala	Ala	Leu 555	Leu	Pro	Ile	Pro	Ala 560
	266																
	266 267	Phe	Arg	Val	Phe	Cys	550 Leu	Gln	Ala	Ala	Ile		Leu	Leu	Phe	Asn	

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/08/916,140

DATE: 10/08/2004 TIME: 18:19:30

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\10082004\H916140.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 4,5,7,8,10,11,23,34,35,36,39,40,41,45,51,52,57,61,71,75,77
Seq#:1; N Pos. 87,88,89,91,92,96,97,100,104,106,109,111,113,117,120,126,149
Seq#:1; N Pos. 151,153,154,157,178,187,189,191,211,214,310,608,704,708,712
Seq#:1; N Pos. 714,729,732
Seq#:2; Xaa Pos. 1,9,12,13,14,20
Seq#:4; Xaa Pos. 348,908

Seq#:7; N Pos. 114,225,261 Seq#:8; Xaa Pos. 75,87

Seq#:14; N Pos. 16,25

Seq#:15; N Pos. 24
Seq#:16; N Pos. 13,16
Seq#:17; N Pos. 20

VERIFICATION SUMMARY

DATE: 10/08/2004 PATENT APPLICATION: US/08/916,140 TIME: 18:19:30

Input Set : A:\PTO.AMC.TXT

Output Set: N:\CRF4\10082004\H916140.raw

L:48 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1 L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0 M:341 Repeated in SeqNo=1 L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 M:341 Repeated in SeqNo=2 L:239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:336 M:341 Repeated in SeqNo=4 L:627 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:60 M:341 Repeated in SeqNo=7 L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:64 M:341 Repeated in SeqNo=8 L:992 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order! L:996 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14 L:997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0 L:1007 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order! L:1011 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15 L:1012 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0 L:1022 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order! L:1026 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16

L:1027 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0

L:1042 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0

L:1037 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order! L:1041 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17



IFW16

RAW SEQUENCE LISTING

DATE: 10/07/2004

PATENT APPLICATION: US/08/916,140

TIME: 10:36:12

Input Set : A:\CIBT-P04-203.TXT

```
4 <110> APPLICANT: Scott Mathew P.
              Goodrich, Lisa V.
              Johnson, Ronald L.
             Epstein, Ervin Jr.
      9 <120> TITLE OF INVENTION: PATCHED GENES AND USES RELATED THERETO
     12 <130> FILE REFERENCE: CIBT-P04-203
     14 <140> CURRENT APPLICATION NUMBER: US 08/916,140
     15 <141> CURRENT FILING DATE: 1997-08-21
     17 <150> PRIOR APPLICATION NUMBER: US 08/656,055
     18 <151> PRIOR FILING DATE: 1996-05-31
     20 <150> PRIOR APPLICATION NUMBER: US 08/540,406
     21 <151> PRIOR FILING DATE: 1995-10-06
     23 <150> PRIOR APPLICATION NUMBER: US 08/319,745
     24 <151> PRIOR FILING DATE: 1994-10-07
     26 <160> NUMBER OF SEQ ID NOS: 64
     28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     30 <210> SEQ ID NO: 1 ·
     31 <211> LENGTH: 736
     32 <212> TYPE: DNA
                                 Beefle
     33 <213> ORGANISM: (Bettle
     35 <220> FEATURE:
     37 <221> NAME/KEY: misc feature
     38 <222> LOCATION: A. 5, 7, 8, 10, 11, /23, 34, /35, 36, 39, 40, 41, 45, 51, 52,
              51, 61, 71, 75, 77, 87, 88, 89, 91, 92, 96, 97, 100, 104, 106, 109, 111, 113, 117, 120, 126, 149, 151, 153, 154, 157, 178, 187, 189, 191, 211, 214, 310, 608, 704, 708, 712
     39
     40
     42 <223> OTHER INFORMATION: n = A, T, C or G
     44 <221> NAME/KEY: misc feature
     45 <222> LOCATION: 7/14, 7/29, 7/32
     46 <223> OTHER INFORMATION: n = A, T, C \text{ or } G
W-->48<400>1
W--> 49 aacnnenntn natggeaece eeneecaaee tttnnneenn ntaancaaaa nneecenttt 60
W--> 50 natacccct ntaananttt tccaccnnnc nnaaannccn ctgnanacna ngnaaanccn 120
W--> 51 tttttnaacc cccccaccc ggaattccna ntnnccnccc ccaaattaca actccagncc 180
W--> 52 aaaattnana naattggtcc taacctaacc natngttgtt acggtttccc cccccaaata 240
     53 catqcactqq cccqaacact tqatcqttqc cqttccaata agaataaatc tggtcatatt 300
W--> 54 aaacaaqccn aaaqctttac aaactgttgt acaattaatg ggcgaacacg aactgttcga 360
     55 attctggtct ggacattaca aagtgcacca catcggatgg aaccaggaga aggccacaac 420
     56 cgtactgaac gcctggcaga agaagttcgc acaggttggt ggttggcgca aggagtagag 480
     57 tgaatggtgg taatttttgg ttgttccagg aggtggatcg tctgacgaag agcaagaagt 540
     58 cgtcgaatta catcttcgtg acgttctcca ccgccaattt gaacaagatg ttgaaggagg 600
W--> 59 cgtcgaanac ggacgtggtg aagctggggg tggtgctggg ggtggcggcg gtgtacgggt 660
W--> 60 gggtggccca gtcggggctg gctgccttgg gagtgctggt cttngcgngc tncnattcgc 720
```

DATE: 10/07/2004

TIME: 10:36:12

Input Set : A:\CIBT-P04-203.TXT Output Set: N:\CRF4\10072004\H916140.raw 736 W--> 61 cctatagtna gncgta 63 <210> SEQ ID NO: 2 64 <211> LENGTH: 107 65 <212> TYPE: PRT 66 <213> ORGANISM: Beetle 68 <220> FEATURE: 70 <221> NAME/KEY: VARIANT 71 <222> LOCATION: 1, 9, 12, 13, 14, 20 72 <223> OTHER INFORMATION: Xaa = Any Amino Acid 74 <400> SEQUENCE: 2 W--> 75 Xaa Pro Pro Pro Asn Tyr Asn Ser Xaa Pro Lys Xaa Xaa Xaa Leu Val W--> 77 Leu Thr Pro Kaa Val Val Thr Val Ser Pro Pro Lys Tyr Met His Trp 78 20 79 Pro Glu His Leu Ile Val Ala Val Pro Ile Arg Ile Asn Leu Val Ile .40 81 Leu Asn Lys Pro Lys Ala Leu Gln Thr Val Val Gln Leu Met Gly Glu 82 83 His Glu Leu Phe Glu Phe Trp Ser Gly His Tyr Lys Val His His Ile 70 75 85 Gly Trp Asn Gln Glu Lys Ala Thr Thr Val Leu Asn Ala Trp Gln Lys 86 85 87 Lys Phe Ala Gln Val Gly Gly Trp Arg Lys Glu 100 91 <210> SEO ID NO: 3 92 <211> LENGTH: 5187 93 <212> TYPE: DNA 94 <213> ORGANISM: Butterfly 96 <400> SEQUENCE: 3 97 gggtctgtca cccggagccg gagtccccgg cggccagcag egtcctcgcg agccgagcgc 60 98 ccaggegege ceggageceg eggeggegge ggeaacatgg ceteggetgg taaegeegee 120 99 ggggccctgg gcaggcaggc cggcggcggg aggcgcagac ggaccgggggg accgcaccgc 180 100 gccgcgccgg accgggacta tctgcaccgg cccagctact gcgacgccgc cttcgctctg 240 101 gagcagattt ccaaggggaa ggctactggc cggaaagcgc cgctgtggct gagagcgaag 300 102 tttcagagac tcttatttaa actgggttgt tacattcaaa agaactgcgg caagtttttg 360 103 gttgtgggtc tcctcatatt tggggccttc gctgtgggat taaaggcagc taatctcgag 420 104 accaacgtgg aggagctgtg ggtggaagtt ggtggacgag tgagtcgaga attaaattat 480 105 acccqtcaga agataggaga agaggctatg tttaatcctc aactcatgat acagactcca 540 106 aaagaagaag gegetaatgt tetgaccaca gaggetetee tgeaacacet ggaeteagea 600 107 ctccaggcca gtcgtgtgca cgtctacatg tataacaggc aatggaagtt ggaacatttg 660 108 tgctacaaat caggggaact tatcacggag acaggttaca tggatcagat aatagaatac 720 109 ctttaccctt gcttaatcat tacacctttg gactgcttct gggaaggggc aaagctacag 780 110 teegggacag cataceteet aggtaageet eetttaeggt ggacaaaett tgaceeettg 840 111 gaatteetag aagagttaaa gaaaataaac taccaagtgg acagetggga ggaaatgetg 900 112 aataaagccg aagttggcca tgggtacatg gaccggcctt gcctcaaccc agccgaccca 960 113 gattgccctg ccacagcccc taacaaaaat tcaaccaaac ctcttgatgt ggcccttgtt 1020 114 ttgaatggtg gatgtcaagg tttatccagg aagtatatgc attggcagga ggagttgatt 1080

115 gtgggtggta ccgtcaagaa tgccactgga aaacttgtca gcgctcacgc cctgcaaacc 1140 116 atgttccagt taatgactcc caagcaaatg tatgaacact tcaggggcta cgactatgtc 1200

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/916,140

RAW SEQUENCE LISTING DATE: 10/07/2004
PATENT APPLICATION: US/08/916,140 TIME: 10:36:12

Input Set : A:\CIBT-P04-203.TXT

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118 tacgtggagg tggttcatca aagtgtcgcc ccaaactcca ctcaaaaggt gcttcccttc 1320
119 acaaccacqa ccctqqacqa catcctaaaa tccttctctg atgtcagtgt catccgagtg 1380
120 gccagcggct acctactgat gcttgcctat gcctgtttaa ccatgctgcg ctgggactgc 1440
121 tocaagtooc agggtgccgt ggggctggct ggcgtcctgt tggttgcgct gtcagtggct 1500
122 gcaggattgg gcctctgctc cttgattggc atttctttta atgctgcgac aactcaggtt 1560
123 ttgccgtttc ttgctcttgg tgttggtgtg gatgatgtct tcctcctggc ccatgcattc 1620
124 aqtqaaacaq qacaqaataa qaqqattcca tttgaggaca ggactgggga gtgcctcaag 1680
125 cgcaccggag ccagcgtggc cctcacctcc atcagcaatg tcaccgcctt cttcatggcc 1740
126 gcattgatcc ctatccctgc cctgcgagcg ttctccctcc aggctgctgt ggtggttggta 1800
127 ttcaattttg ctatggttct gctcattttt cctgcaattc tcagcatgga tttatacaga 1860
128 cgtgaggaca gaagattgga tattttctgc tgtttcacaa gcccctgtgt cagcagggtg 1920
129 attcaagttg agccacaggc ctacacagag cctcacagta acacceggta cagcccccca 1980
130 cccccataca ccagccacag cttcgcccac gaaacccata tcactatgca gtccaccgtt 2040
131 cageteegea cagagtatga ceeteacaeg caegtgtaet adaceaeege egageeaege 2100
132 tetqaqatet etqtacagee tgttacegte acceaggaca accteagetg teagagteee 2160
133 gagageacca getetaccag ggacetgete teccagttet cagactecag cetecactge 2220
134 ctcqaqcccc cctqcaccaa qtggacactc tcttcgtttg cagagaagca ctatgctcct 2280
135 ttcctcctga aacccaaagc caaggttgtg gtaatccttc ttttcctggg cttgctgggg 2340
136 gtcagccttt atgggaccac ccgagtgaga gacgggctgg acctcacgga cattgttccc 2400
137 cgggaaacca gagaatatga cttcatagct gcccagttca agtacttctc tttctacaac 2460
138 atgtatatag teacceagaa ageagaetae eegaatatee ageacetaet ttaegaeett 2520
139 cataagagtt tcagcaatgt gaagtatgtc atgctggagg agaacaagca acttccccaa 2580
140 atqtqqctqc actactttaq aqactqqctt caaggacttc aggatgcatt tgacagtgac 2640
141 tqqqaaactq qqaqqatcat qccaaacaat tataaaaatg gatcagatga cggggtcctc 2700
142 gcttacaaac teetggtgea gaetggeage egagacaage ceategaeat tagteagttg 2760
143 actaaacage gtetggtaga egeagatgge atcattaate egagegettt etacatetae 2820
144 etgacegett gggteageaa egaceetgta gettaegetg eeteceagge caacateegg 2880
145 cctcaccggc cggagtgggt ccatgacaaa gccgactaca tgccagagac caggctgaga 2940
146 atcccagcag cagageccat cgagtacget cagtteeett tetaceteaa eggeetacga 3000
147 gacacctcag actttgtgga agccatagaa aaagtgagag tcatctgtaa caactatacg 3060
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149 agectgegee actggetget getatecate agegtggtge tggeetgeae gtttetagtg 3180
150 tgcgcagtct tcctcctgaa cccctggacg gccgggatca ttgtcatggt cctggctctg 3240
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155 tecgaatttg attteattgt cagatactte tttgeegtee tggeeattet caeegtettg 3540
156 ggggttetea atggaetggt tetgetgeet gteetettat eettetttgg accgtgteet 3600
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158 agtgtcgtcc ggtttgccgt gcctcctggt cacacgaaca atgggtctga ttcctccgac 3720
159 teggagtaca geteteagae caeggtgtet ggeateagtg aggageteag geaataegaa 3780
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163 cgaagggate eccetagaga aggettgegg ceaececeet acagacegeg cagagaeget 4020
164 tttgaaattt ctactgaagg gcattctggc cctagcaata gggaccgctc agggccccgt 4080
165 qqqqcccqtt ctcacaaccc tcqqaaccca acgtccaccg ccatgggcag ctctgtgccc 4140
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RAW SEQUENCE LISTING DATE: 10/07/2004
PATENT APPLICATION: US/08/916,140 TIME: 10:36:12

Input Set : A:\CIBT-P04-203.TXT

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168 cctgagactg atcacggggt atttgaggat cctcatgtgc cttttcatgt caggtgtgag 4320
169 aqqaqqact caaaqqtqqa qqtcataqaq ctacaqqacg tggaatgtga ggagaggccg 4380
170 tgggggagca getecaactg agggtaatta aaatetgaag caaagaggee aaagattgga 4440
171 aagccccgcc cccacctctt tccagaactg cttgaagaga actgcttgga attatgggaa 4500
172 ggcagttcat tgttactgta actgattgta ttattkkgtg aaatatttct ataaatattt 4560
173 aaraggtgta cacatgtaat atacatggaa atgctgtaca gtctatttcc tggggcctct 4620
174 ccactcctgc cccagagtgg ggagaccaca ggggcccttt cccctgtgta cattggtctc 4680
175 tgtgccacaa ccaagcttaa cttagtttta aaaaaaatct cccagcatat gtcgctgctg 4740
176 ottaaatatt gtataattta ottgtataat totatgoaaa tattgottat gtaataggat 4800
177 tatttgtaaa ggtttctgtt taaaatattt taaatttgca tatcacaacc ctgtggtagg 4860
178 atgaattgtt actgttaact tttgaacacg ctatgcgtgg taattgttta acgagcagac 4920
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181 tgctgttgtt gttcattttg gtgtttttgg ttgctttgta tgatcttagc tctggcctag 5100
182 gtgggctggg aaggtccagg tctttttctg tcgtgatgct ggtggaaagg tgaccccaat 5160
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185 <210> SEQ ID NO: 4
186 <211> LENGTH: 1311
187 <212> TYPE: PRT
188 <213> ORGANISM: Butterfly
190 <220> FEATURE:
192 <221> NAME/KEY: VARIANT
193 <222> LOCATION: 348, 908
194 <223> OTHER INFORMATION: Xaa = Any Amino Acid
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198
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200
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201 Tyr Ile Arg Thr Ser Trp Val Asp Ala Ala Leu Ala Leu Ser Glu Leu
203 Glu Lys Gly Asn Ile Glu Gly Gly Arg Thr Ser Leu Trp Ile Arg Ala
204
                            55
205 Trp Leu Gln Glu Gln Leu Phe Ile Leu Gly Cys Phe Leu Gln Gly Asp
206 65
207 Ala Gly Lys Val Leu Phe Val Ala Ile Leu Val Leu Ser Thr Phe Cys
208
                                        90
                    85
209 Val Gly Leu Lys Ser Ala Gln Ile His Thr Arg Val Asp Gln Leu Trp
                                    105
211 Val Gln Glu Gly Gly Arg Leu Glu Ala Glu Leu Lys Tyr Thr Ala Gln
                                                    125
                                120
212
213 Ala Leu Gly Glu Ala Asp Ser Ser Thr His Gln Leu Val Ile Gln Thr
       130
                            135
215 Ala Lys Asp Pro Asp Val Ser Leu Leu His Pro Gly Ala Leu Leu Glu
                        150
                                            155
217 His Leu Lys Val Val His Ala Ala Thr Arg Val Thr Val His Met Tyr
                    165
                                        170
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/916,140

DATE: 10/07/2004 TIME: 10:36:12

Input Set : A:\CIBT-P04-203.TXT

		Asp	Ile	Glu		Arg	Leu	Lys	Asp		Cys	Tyr	Ser	Pro		Ile	Pro
	220 221	Asp	Phe	Ġlu	180 Gly	Tyr	His	His		185 Glu	Ser	Ile	Ile	Asp	190 Asn	Val	Ile
	222	Dro	Crra	195	T10	Tlo	Thr	Dro	200	7) cm	Cys	Dhe	Trn	205 Glu	Glv	Ser	Lve
	224		210					215					220				
	225 226		Leu	Gly	Pro	Asp	Tyr 230	Pro	Ile	Tyr	Val	Pro 235	His	Leu	Lys	His	Lys 240
			Gln	Trp	Thr	His		Asn	Pro	Leu	Glu		Val	Glu	Glu		
	228	T	T 011	Tira	Dho	245	Dho	Dro	Lou	cor	250 Thr	Tla	Glu	Λla	Tur	255 Met	Larg
	230	гуѕ	ьеи	пуъ	260	GIII	FILE	FIO	пеп	265	1111	110	Oiu	2114	270	rice	цуб
		Arg	Ala	Gly		Thr	Ser	Ala	Tyr	Met	Lys	Lys	Pro	Cys	Leu	Asp	Pro
	232			275				_ =	280		_	_	_	285		~ 1	·
		Thr	_	Pro	His	Cys	Pro	Ala 295	Thr	Ala	Pro	Asn	300	ьуs	ser	GIY	HIS
	234	Tle	290 Pro	Asp	Val	Ala	Ala		Leu	Ser	His	Glv		Tvr	Glv	Phe	Ala
	236				• 0		310					315			_		320
	237	Ala	Ala	Tyr	Met	His	Trp	Pro	Glu	Gln	Leu	Ile	Val	Gly	Gly	Ala	Thr
	238					325					330					335	
W>		Arg	Asn	Ser		Ser	Ala	Leu	Arg		Ala	Arg	Xaa	Leu	Gln 350	Thr	Val
	240	17a l	Cln	Lou	340	Clar	Glu	Λrα	Glu	345 Met	Tyr	Glu	Tur	Tro		Asp	His
	241	vai	GIII	355·	MEC	СТУ	Giu	Arg	360	MCC	1 y 1.	GIU	- y -	365	1114	тър	113.6
		Tvr	Lys		His	Gln	Ile	Gly		Asn	Gln	Glu	Lys	Ala	Ala	Ala	Val
	244		370					375					380				
			Asp	Ala	Trp	Gln		Lys	Phe	Ala	Ala		Val	Arg	Lys	Ile	
		385	_		_	7	390	_		_		395	m. i	D	D1	0	400
	247 248	Thr	Ser	GIY	ser	Val 405	ser	ser	Ата	Tyr	Ser 410	Pne		PIO		415	TIII
		Ser	Thr	T.eu	Asn		Tle	Leu	Glv	Lvs	Phe	Ser					Lvs
	250	DCL		Lea	420	тыр		200	0-1	425					430		_
	251	Asn	Ile	Ile	Leu	Gly	Tyr	Met	Phe	Met	Leu	Ile	Tyr	Val	Ala	Val	Thr
•	252	_		435		_	_	_	440	_	G	α1	n 7 -	445	TT 7	al. .	T] a:
	253		11e 450	GIn	Trp	Arg	Asp	455	TTE	arg	Ser	GIII	460	GIY	Val	GIY	11e
			Gly	Val	Leu	Leu	Leu	Ser	Ile	Thr	Val	Ala	Ala	Gly	Leu	Gly	Phe
	256	465					470					475					480
			Ala	Leu	Leu		Ile	Pro	Phe	Asn	Ala	Ser	Ser	Thr	Gln		Val
	258		Db.a	T	7. 7	485	~ 1	T 011	C1	Tro 1	490	7 an	Mot	Dho	Lau	495	Thr
	260	PIO	Pile	ьeu	500	ьеи	GIY	ьеи	СТУ	505	GIII	Asp	Mec	FILE	510	i i i	Thr
		His	Thr	Tyr		Glu	Gln	Ala	Gly		Val	Pro	Arg	Glu		Arg	Thr
	262			515					520	_			_	525			
	263	Gly	Leu	Val	Leu	Lys	Lys	Ser	Gly	Leu	Ser	Val	Leu	Leu	Ala	Ser	Leu
	264		530			-		535				_	540				2.7
			Asn	Val	Met	Ala		Leu	Ala	Ala	Ala		Leu	Pro	тте	Pro	A1a 560
		545	Δ×~	v-1	Dhe	Cvc	550	Gln	د 1 ۵	د [۵	Ile	555 Leu	Leu	Len	Phe	Asn	
	201	rne	ALY	vaı	FIIG	Cys	пеи	GIII	vra	AIG	116	±-cu		Leu	1110	11011	

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 10/07/2004

PATENT APPLICATION: US/08/916,140

TIME: 10:36:13

Input Set : A:\CIBT-P04-203.TXT

Output Set: N:\CRF4\10072004\H916140.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. A,8,7,8,10,11,23,34,35,36,39,40,41,45,51,52,57,61,71,75,77, Seq#:1; N Pos. 87,88,89,91,92,96,97,100,104,106,109,111,113,117,120,126,149 Seq#:1; N Pos. 151,153,154,157,178,187,189,191,211,214,310,608,704,708,712 Seq#:1; N Pos. 714,729,732

Seq#:2; Xaa Pos. 1,9,12,13,14,20

Seq#:4; Xaa Pos. 348,908

Seq#:7; N Pos. 114,225,261

Seq#:8; Xaa Pos. 75,87

Seq#:14; N Pos. 16,25

Seq#:15; N Pos. 24 Seq#:16; N Pos. 13,16

Seq#:17; N Pos. 20

VERIFICATION SUMMARY

PATENT APPLICATION: US/08/916,140

DATE: 10/07/2004 TIME: 10:36:13

Input Set : A:\CIBT-P04-203.TXT

```
L:48 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:60
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:120
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:300
L:59 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:600
L:60 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:660
L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:720
L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:16
L:239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:336
L:309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:896
L:627 \ M:341 \ W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:60
L:629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:180
L:630 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:240
L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:64
L:655 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:80
L:992 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:996 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:1007 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:1011 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:1012 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:1022 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:1026 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:1027 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:1037 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:1041 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:1042 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
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